

# SEQUENCE LISTING

<110> Nycomed Pharma AS  
 Ihle, Oystein  
 Michaelsen, Terje Einar  
 Johnne, Berit  
 Grant, Anne R

<120> Method

<130> 28.69235/001

<140> PCT/GB99/03995

<141> 1999-11-30

<150> GB9826247.0

<151> 1998-11-30

<160> 26

<170> PatentIn Ver. 2.1

<210> 1

<211> 15

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Target fragment  
 for Bam HI primer

<400> 1

tttactggat cctag

15

<210> 2

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: target  
 fragment for Asn I primer

<400> 2

ttacgtacat taatcgg

17

<210> 3

<211> 17

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Asn I primer

<400> 3

ccgattaatg tacgtaa

17

<210> 4

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Incorrect  
restriction site in fragment

<400> 4

ttacgtacaa tcgg

14

<210> 5

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Incorrect  
restriction site in fragment

<400> 5

tttactggta g

11

<210> 6

<211> 12

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Incorrect  
restriction site in fragment

<400> 6

tttactggat ag

12

<210> 7

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:)Correct  
restriction site in fragment (complementary  
strand)

<400> 7

gatccagtaa a

11

<210> 8

<211> 13

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Incorrect  
restriction fragment (complementary strand)

<400> 8

ttaatgtacg taa

13

<210> 9

<211> 11

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Incorrect  
restriction fragment (complementary strand)

<400> 9

ctaccagtaa a

11

<210> 10

<211> 14

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Incorrect  
restriction fragment (complementary strand)

<400> 10

ccgattgtac gtaa

14

<210> 11

<211> 17

09870128 "053004  
T00000

<212> DNA  
<213> Homo sapiens

<400> 11  
atgacctagg accacct 17

<210> 12  
<211> 17  
<212> DNA  
<213> Homo sapiens

<400> 12  
atgacctagg cccacct 17

<210> 13  
<211> 93  
<212> DNA  
<213> Homo sapiens

<400> 13  
atggtgcacc tgactcctga ggagaagtct gccgttactg ccctgtgggg caaggtgaac 60  
gtggatgaag ttggtggtga ggccctgggc agg 93

<210> 14  
<211> 30  
<212> PRT  
<213> Homo sapiens

<400> 14  
Met Val His Leu Thr Pro Glu Glu Lys Ser Ala Val Thr Ala Leu Trp  
1 5 10 15

Gly Lys Val Asn Val Asp Glu Val Gly Gly Glu Ala Leu Gly  
20 25 30

<210> 15  
<211> 93  
<212> DNA  
<213> Homo sapiens

<400> 15  
atggtgcacc tgactcctgt ggagaagtct gccgttactg ccctgtgggg caaggtgaac 60  
gtggatgaag ttggtggtga ggccctgggc agg 93

<210> 16  
<211> 30  
<212> PRT

<213> Homo sapiens

<400> 16

Met Val His Leu Thr Pro Val Glu Lys Ser Ala Val Thr Ala Leu Trp  
1 5 10 15

Gly Lys Val Asn Val Asp Glu Val Gly Gly Glu Ala Leu Gly  
20 25 30

<210> 17

<211> 306

<212> DNA

<213> Homo sapiens

<400> 17

gcataaaagt cagggcagag ccattctattg cttacatttg cttctgacac aactgtgttc 60  
actagcaacc tcaaacagac accatggtgc acctgactcc tgaggagaag tctgccgtta 120  
ctgccctgtg gggcaaggtg aacgtggatg aagttggtgg tgaggccctg ggcaggggca 180  
ggttggtatc aaggttataa gacaggttta aggagaccaa tagaaactgg gcatgtggag 240  
acagagaaga ctcttggtt tctgataggc actgactctc tctgcctatt ggtctatttt 300  
cccacc 306

<210> 18

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 18

ctagcaacct caaacagaca cc 22

<210> 19

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 19

gtaaccttga taccaacctg cc 22

<210> 20

<211> 20

<212> DNA

09870128-053004

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 20

atggtgcacc tgactcctga

20

<210> 21

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 21

atggtgcacc tgactcctgt

20

<210> 22

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 22

gtaaccttga taccaacctg cc

22

<210> 23

<211> 93

<212> DNA

<213> Homo sapiens

<400> 23

atggtgcacc tgactcctaa cgagaagtct gccgttactg ccctgtgggg caaggtgaac 60

gtggatgaag ttggtggtga ggccctgggc agg

93

<210> 24

<211> 30

<212> PRT

<213> Homo sapiens

<400> 24

Met Val His Leu Thr Pro Asn Glu Lys Ser Ala Val Thr Ala Leu Trp

1

5

10

15

Gly Lys Val Asn Val Asp Glu Val Gly Gly Glu Ala Leu Gly  
 20 25 30

<210> 25

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 25

atggtgcacc tgactcctaa c 21

<210> 26

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 26

atggtgcacc tgactccta 19

09870128-053004